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Box Missing Parts, Washington, D.C. 20231, on

11-26-01

LAW OFFICES OF JONATHAN ALAN QUINE

Attorney Docket No. 02-106720US

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re application of:

Juha Punnonen, et al.

Application No.: 09/888,324

Filed: June 22, 2001

NOVEL CO-STIMULATORY

MOLECULES

Examiner: Unassigned

Art Unit: 1643

LETTER TO OFFICIAL DRAFTSPERSON

Attn:

For:

Assistant Commissioner for Patents

Washington, D.C. 20231

Sir:

Applicant hereby submits 39 sheets of formal drawings to be made of record in the above-identified case.

Respectfully submitted,

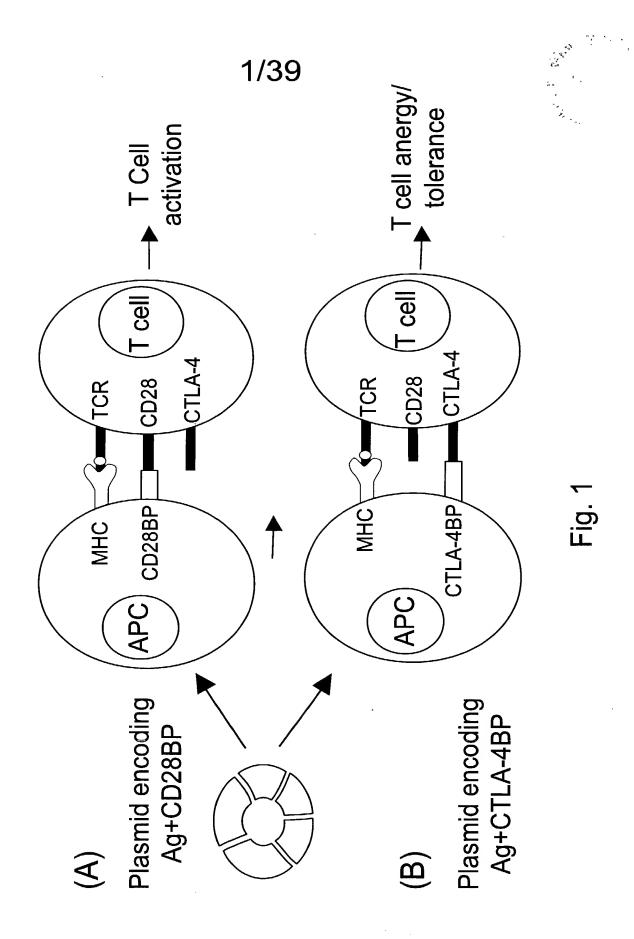
Emily M. Haliday, J.D., Ph. Reg. No. 38,903

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2 0



MGHTRRQGTSPSKCPYLNFFQLLVLAGLSHFCSG--VIHVTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMS Signal sequence

(1)

MGHTLRPGTPLPRCLHLKLCLLLALAGLHFSSG---ISQVTKSVKEMAALSCDYNISIDELARMRIYWQKDQQMVLSIIS MGHTLRPGTPL PRCLHLKLCLLLALAGLHFSSG---ISQVTKSVKEMAALSCDYNISIDELARMRIYWQKDQQMVLSIIS MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKMVLA1LP (1)

> SEQ:050_R1_Clone_118 SEQ:051_R1_Clone_126

SEQ:048_R1_Clone_71 SEQ:049_R1_Clone_84 SEQ:052_R2_CD28BP-1 SEQ:053_R2_CD28BP-2 SEQ:054_R2_CD28BP-3 SEQ:055_R2_CD28BP-4 SEQ:056_R2_CD28BP-5

SEQ:278_Human_B7-1

MGHTWKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYSTSTEELTSLR1VWQKDSKMVLA1LP (1)

MGHTWKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP (1)

MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP

MGHTMKWGSL PPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAIĻP <u>1</u>

MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP (1)

MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTRRVKETVMLSCDYSTSTEELTSLRIYWOKDSKMVLAILP (1)

MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKMVLAILP 1 (1)

MGHTMKWRSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP 1

MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWRSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELJSLRIYWQKDSKMVLAILP (1) (1)

SEQ:061_R2_CD28BP-10

SEQ:062_R2_CD28BP-11 SEQ:063_R2_CD28BP-12 SEQ:064_R2_CD28BP-13 SEQ:065_R2_CD28BP-14

SEQ:059_R2_CD28BP-8 SEQ:060_R2_CD28BP-9

SEQ:057_R2_CD28BP-6

SEQ:058_R2_CD28BP-7

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MGHTLRPGTPLPRCLHLKLCLLLALAGLHFSSG---ISQVTKSVKEMAALSCDYNISIDELARMRIYWQKDQQMVLSIIS (1)

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MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEKLTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWRKDSKMXLAILP MGHTMKWGSLPPKCPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP (1) <u>1</u> (1) SEQ:174_cd28A12-5 SEQ:175_cd28a4-5star

MGHTWKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKMVLA1LP MGHTWKWGSLPPKRPCLWLPQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP 1 (1) SEQ:176_cd28A4-9 SEQ:177_cd28A6-9 SEQ:178_cd28A6-1

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MGHTMKWGSL PPKR PCLWL SQLLVL TGL FY PCSGITPKSVTKRVK BTVML SCDYNTSTEEL TSLR I YWQKDSKMVLAIL P MGHTWKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKWYLAILP (1) (1) SEQ:181_cd28B2-8 SEQ:182_cd28B4-3

MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKMVLAILP MGHIMKNGSL PPKR PCLWLSQLLVL TGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEEL TSLR IYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLR1YWQKDSKMVLAILP (1) SEQ:184_cd28b6-6 SEQ:185_cd28b8-5star SEQ:183_cd28B6-3

MGHTMKWGSLPPKRPCLWPSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKMVLAILP

SEQ:186_cd28c11-5



Extracellular domain (ECD)

Signal sequence -

MGHTWKWGSLPPKRPCLWLSOLLVLTGLFYFCSG1TPKSVTKRVKFTVMLSCDYSTSTEELTSLR1YWOKDSKMYLA1LP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYSTSTEELTSLR1YWQKDSKMVLA1LP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKMVLA1LP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLR1YWQKDSKMVLAILP MGHTWKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGTTPKSVTKRVKETVMLSCDYSTSTEELTSLR1YWQXDSKMVLAILP MGHTMEWGSL PPKR PCLWL SQLLVL TGL FYFCSGI TPKSVTKRVK ETVML SCDYNTSTEEL TSLRI YWQKDSKMVLAIL P MGHTMKWGSL PPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQXDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKMVLAILP MGHTMKWGSL PPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYSTSTEELTSLR1YWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKMVLA1LP MGHTMKWGSL PPKRPCLWLSQLLVLTDLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKMYLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTRRVKETVMLSCDYSTSTEELTSLRIYWOKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYSTSTEELTSLR1YWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKMVLA1LP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKMVLA1LP MGHTMKWGSL PPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKMVLA1LP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSGTTPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKMVLAILP MGHTMKWGSL PPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP MGHTWKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKMYLAILP MGHTWKWGSLPPKRPCLMLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELJSLRIYWQKDSKMVLAILP MGHTWKWGSLPPKRPCLRLSQLLVLTGLFYFCSG1TPKSVTKRVXETVMLSCDYSTSTEELTSLR1YWQKDSKMYLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKMVLAILP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYFCSG1TPKSVTKRVKETVMLSCDYNTSTEELTSLR1YWQKDSKMYLAILP MGHTWKWGSLPPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYSTSTEELTSLRIYWQXDSKWVLAILP MGHTLR PGTPL PRCLHLKLCLLLALAGLHFSSG---ISQVTKSVKEMAALSCDYNISIDELARMRIYWOKDQOMVLSIIS MGHTMKWGSLPPRRPCLWLSQLLVLTGLFYFCSG1TPRSVTKRVKETVMLSCDYNTSTEELTSLR1YWOXDSKMVLA1LP MGHTMKWGSLPPKRPCLWLSQLLVLTGLFYPCSG1TPKSVTKRVKETVMLSCDYSTSTEELTSLR1YWQXDSKMVLA1LP MGHTMKWGSL PPKRPCLWLSQLLVLTGLFYFCSGITPKSVTKRVKETVMLSCDYNTSTEELTSLRIYWQKDSKMVLAILP (1)(1) (1) $\widehat{\mathbf{1}}$ (1) (T $\widehat{\mathbf{1}}$ (1) (1) 1 (1)(1)(1) $\widehat{\Xi}$ $\widehat{\Xi}$ $\widehat{\mathbf{1}}$ 1) (1) $\widehat{\Xi}$ $\widehat{\Xi}$ (1) (1) 1 $\widehat{\exists}$ $\widehat{\mathbf{1}}$ $\widehat{\mathbf{I}}$ SEQ:210_cd28A2-5a SEQ:191_cd28C2-4 SEQ:192_cd28D2-3 SEQ:193_cd28D2-9 SEQ:194_cd28D8-9 SEQ:196_cd28D12-5 SEQ:197_cd28E10-6 SEQ:198_cd28F7-2 SEQ:199_cd28F8-4 SEQ:200_cd28F10-2 SEQ:201_cd28F12-5star SEQ:202_cd28G2-8 SEQ:203_cd28G1-5 SEQ:204_cd28G1-9 SEQ:205_cd28H4-3 SEQ:206_cd28H11-3 SEQ:207_cd28H6-6 SEQ:209_cd28B4-5a SEQ:211_cd28B4-5star SEQ:212_cd28D5-6 SEQ:213_cd28D10-4 SEQ:214_cd28E2-5star SEQ:215_cd28E5-2 SEQ:216_cd28E8-6 SEQ:217_cd28E9-6 SEQ:219_cd28F3-5 SEQ:221_cd28F11-8 SEQ:187_cd28C6-1 SEQ:188_cd28C7-3 SEQ:189_cd28C8-6 SEQ:190_cd28c9-5star SEQ:195_cd28D11-1 SEQ:208_cd28E2-4 SEQ:218_cd28F3-1 SEQ:220_cd28F3-6

Fig. 2B

GDMNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLK-YEKDAFKREHLAEVTLSVKADFPTPSISDFEIPTSNI GKVQVWPEYKNRTITDMADNPRIVILALRLSDKGTYTCVVQKPVLKGAYKLEHLTSVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNLRIVILALRLSDSGTYTCVIQKPDLKGAYKLEHLTSVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVVQK-NENGSFRREHLTSVTLSIRADFPVPSITDIGHPAPNV GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPDLKGAYKLEHLTSVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDKGTYTCVIQKPVLKGAYKLEHLTSVTLSIRADFPVPSITDIGHPAPNV GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI GQVEVWPEYKNRTFPDIINNLSLMILALRLSDKGTYTCVVQK-NENGSFRREHLTSVTLSIRADSPVPSITDIGHPAPNV GQVEVWPEYKNRTFPDIINNLSLMILALRLSDKGTYTCVVQK-NENGSFRREHLTSVTLSIRADFPVPSITDIGHPAPNV GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPDLKGAYKLEHLTSVRLMIRADFPVFTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVFTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPSITDIGHPAPNV GKVQVWPEYKNRTITDMADNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLTSVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMADNPRIVILALRLSDKGTYTCVVQK-NENGSFRREHLTSVTLSIRADFPVPSITDIGHPAPNV GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI GQVEVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRPSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTFPDIINNLSLMILALRLSDKGTYTCVVQK-NENGSFRREHLTSVTLSIRADFPVPSITDIGHPAPNV GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLTSVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLTSVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDKGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFFVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLTSVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLTSVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLTSVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLTSVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLTSVRLMIRADFPVPTINDLGNPSPNI Extracellular domain (ECD) (42) 81) 81) (81) (81) (18) (81) (81) 81) 81) 81) (81) 81) 81) (81) 81) 81) (81)(81)81) 81) (81) 81) 81) 81) 81) 81) 81) 81) (81)81) 78) SEQ:176_cd28A4-9 SEQ:278_Human_B7-1 SEQ:049_R1_Clone_84 SEQ:050_R1_Clone_118 SEQ:051_R1_Clone_126 SEQ:052_R2_CD28BP-1 SEQ:053_R2_CD28BP-2 SEQ:054_R2_CD28BP-3 SEQ:055_R2_CD28BP-4 SEQ:056_R2_CD28BP-5 SEQ:057_R2_CD28BP-6 SEQ:058_R2_CD28BP-7 SEQ:059_R2_CD28BP-8 SEQ:060_R2_CD28BP-9 SEQ:061_R2_CD28BP-10 SEQ:062_R2_CD28BP-11 SEQ:063_R2_CD28BP-12 SEQ:064_R2_CD28BP-13 SEQ:065_R2_CD28BP-14 SEQ:066_R2_CD28BP-15 SEQ:067_R2_CD28BP-16 SEQ:068_R2_CD28BP-17 SEQ:174_cd28A12-5 SEQ:175_cd28a4-5star SEQ:177_cd28A6-9 SEQ:178_cd28A6-1 SEQ:179_cd28A8-4 SEQ:180_cd28A8-6 SEQ:181_cd28B2-8 SEQ:182_cd28B4-3 SEQ:183_cd28B6-3 SEQ:184_cd28b6-6 SEQ:185_cd28b8-5star SEQ:186_cd28c11-5 SEQ:048_R1_Clone_71

Fig. 20



160

GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPDLKGAYKLEHLTSVRLMIRADFPVPSITDIGHPAPNV

GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI

81

(81) (81)

SEQ:187_cd28C6-1 SEQ:188_cd28C7-3

(81)

SEQ: 189_cd28C8-6

SEQ:190_cd28c9-5star SEQ:191_cd28C2-4

(81) (42)

(81) (81)

SEQ:192_cd28D2-3 SEQ:193_cd28D2-9

Extracellular domain (ECD)

GKVQVWPEYKNRTITDMMDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI GQVEVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKPEHLASVRLMIRADFPVPTINDLGNPSPNI GKVQVWPEYKNRTITDMNDNPRIVIQALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVP--TDLGNPSPNI GKVQVWPEYKNRTITIDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLTSVRLMIRADFPVPTINDLGNPSPNI

,, ,,	(81) GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLTSVRLMIRADFPVPTINDLGNPSPNI (81) GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIOKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI
· .;	
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	(81) GKVQVWPEYKNRTITDMADNPRIVILALRLSDKGTYTCVIQKPDLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI
	(81) GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLTSVRLMIRADFPVPTINDLGNPSPNI
	(81) GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPDLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI
	(81) GKVQVWPEYKNRTITDMADNPRIVILALRPSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI
	(81) GKVQVWPEYKNRTITDMNDNPRIVILALRLSDKGTYTCVVQK-NENGSFRREHLTSVTLSIRADFPVPSITDIGHPAPNV
	(81) GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLTSVRLMIRADFPVPTINDLGNPSPNI
	(81) GKVQVWPEYKNRTITDMADNPRIVILALRLSDKGTYTCVVQKPDLKGAYKLEHLASVRLMIRADFPVPSITDIGHPAPNV
	(81) GKVQVWPEYKNRTITDMADNPRIVILALRLSDSGTYTCVIQKPDLKGAYKLEHLASVRLMIRADFPVPSITDIGHPAPNV
	(81) GKVQVWPEYKNRTITDNMDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI
	(81) GKVQVWPEYKNRTITDMNDNPRIVILALRLSDKGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI
1	(81) GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI
) ((81) GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI
(3	(81) GKVQVWPEYKNRTIITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLTSVRLMIRADFPVPSITDIGHPAPNV
)/	(81) GKVQVWPEYKNRTFPDIINNLSLMILALRLSDKGTYTCVVQK-NENGSFRREHLTSVTLSIRADFPVSSITDIGHPAPNV
	(81) GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPSINDLGNPSPNI
	(81) GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPDLKGAYKLEHLASVRLMIRADFPVPSITDIGHPAPNV
	(81) GKVQVWPEYKNRTITDMMDNPRIVILALRLSDSGTYTCVIQKPDLKGAYKLEHLTSVRLMIRADFPVPSITDIGHPAPNV
	(81) GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI
	(81) GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI
	(81) GKVQVWPEYKNRTITDMMDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADFPVPTINDLGNPSPNI
	(81) GKVQVWPEYKNRTITDMADNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLASVRLMIRADPPVPSITDIGHPAPNV
	(81) GKVQVWPEYKNRTITDMNDNPRIVILALRLSDSGTYTCVVQK-NENGSFRREHLTSVTLSIRADFPVPSITDIGHPAPNV
	(81) GKVQVWPBYKNRTITDMNDNPRIVILALRLSDSGTYTCVIQKPVLKGAYKLEHLTSVRLMIRADFPVPTINDLGNPSPNI

RRLICSTSGGFPRPHLYWLENGEELNATNTTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATNTTLSQDPETKLYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATNTTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATNTTVSQDPETKLYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATNTTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATNTTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE KRIRCSASGGFPEPRLAMMEDGEELNAVNTTVDQDLDTELYSVSSELDFNATNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATWTTLSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE (158) RRIICSTSGGFPEPHLSWLENGEELNAINTTVSQDPETELYAVSSKLDFNMTTNHSFMCLIKYGHLRVNQTFNWNTTKQE KRIRCSASGGFPEPRLAWMEDGEELNAVNTTVDQDLDTELYSVSSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE KRIRCSASGDFPEPRLAWMEDGEELNAVNTTVDQDLDTELYSVSSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATUTUSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATNTTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLCWLENGEELNATNTTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE rrlicstsggfprphlywlengeelnatnttvsqdpgtelymisseldfnvtnnhsivclikygelsvsqifpwskpkqe RRLICSTSGGFPRPHLYWLENGEELNATNTTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE KRIRCSASGGFPEPRLAWMEDGEELNAVNTTVDQDLDTELYSVSSELDSNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATNTTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE rrlicstsggfprphlywlengeelnatnttlsqdpgtelymisseldfnytnnhsivclikygelsvsqifpwskpkqe RRLICSTSGGFPEPRLAMMEDGEELNAVNTTVDQDLDTELYSVSSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATNTTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE KRIRCSASGDFPEPRLAMMEDGEELNAVNTTVDQDLDTELYSVSSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATNTTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELLVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATNTTLSQDPETKLYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATNTTVSQDPETKLYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATNTTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE KRIRCSASGGFPEPRLAMMEDGEELNAVNTTVDQDLDTELYSVSSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRL I CSTSGGFPR PHL YWL ENGEEL NA TNTTL SQDPETKL YM I SSEL DFNMTSNHSFLCL VKYGDL TV SQTFYWQESK PT RRLICSTSGGFPRPHLYWLENGEELNATNTTLSQDPETKLYMISSELDFNMTSNHSFLCLVKYGDLTVSQTFYWQESKPT RRLICSTSGGFPRPHLYWLENGEELNATNTVSQDPGTELYMISSELDFNVTNNHSIACLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATUTTLSQDPETKLYMISSELDFNMTSNHSFLCLVKYGDLTVSQTFYWQESKPT KRIRCSASGGFPEPRLAWMEDGEELNAVNTTVDQDLDTELYSVSSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATNTTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATNTTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATNTTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKOE 161 (157)(161)(161)(157)(161)(191) (161)(161)(161)(161)(161)(161)(161)(161)(161)(160)(161)(161)(158)(161)(161)(160)(161)(161)(161)(161)(161)(160)(161)(161)(161)(161)(161)(161)SEQ:051_R1_Clone_126 SEQ:067_R2_CD28BP-16 SEQ:174_cd28A12-5 SEQ:175_cd28a4-5star SEQ:176_cd28A4-9 SEQ:177_cd28A6-9 SEQ:178_cd28A6-1 SEQ:181_cd28B2-8 SEQ:183_cd28B6-3 SEQ:186_cd28c11-5 SEQ:278_Human_B7-1 SEQ:050_R1_Clone_118 SEQ:053_R2_CD28BP-2 SEQ:054_R2_CD28BP-3 SEQ:055_R2_CD28BP-4 SEQ:056_R2_CD28BP-5 SEQ:057_R2_CD28BP-6 SEQ:059_R2_CD28BP-8 SEQ:060_R2_CD28BP-9 SEQ:061_R2_CD28BP-10 SEQ:063_R2_CD28BP-12 SEQ:064_R2_CD28BP-13 SEQ:065_R2_CD28BP-14 SEQ:066_R2_CD28BP-15 SEQ:068_R2_CD28BP-17 SEQ:179_cd28A8-4 SEQ:180_cd28A8-6 SEQ:182_cd28B4-3 SEQ:184_cd28b6-6 SEQ:185_cd28b8-5star SEQ:048_R1_Clone_71 SEQ:049_R1_Clone_84 SEQ:052_R2_CD28BP-1 SEQ:058_R2_CD28BP-7 SEQ:062_R2_CD28BP-11

KRIRCSASGDFPEPRLAMMEDGEELNAVNTTV---LDTELYSVSSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPEPRLAWMEDGEELNAVNTTVDQDLDTELYSVSSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE KRIRCSASGDFPEPRLAWMEDGEELNAVNTTVDQDLDTELYSVSSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICS---GFPRPHLYWLENGEELNATUTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATUTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELMATUTTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE KR IRCSASGGFPEPRLAWMEDGEELNAVNTTVDQDLDTELYSVSSELDFNMTSNHSFLCLVKYGDLTVSQTFYWQESKPT KRIRCSASGGFPEPRLAWMEDGEELMAVNTTVDQDLDTELYSVSSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE KRIRCSASGGFPEPRLAMMEDGEELNAVNTTVDQDLDTELYSVSSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE KRIRCSASGGFPEPRLAWMEDGEELNAVNTTVDQDLDTELYSVSSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE rrlicstsggfprphlywlengeelnatntvsqdpdtelymisseldfnvtnnhsivclkygelsvsqifpwskpkqe RRLICSTSGGFPRPHLYWLENGEELNATUTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATNTTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATUTTLSQDPFTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATNTTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE KRIRCSASGDFPEPRLAWMEDGEELNAVNTTVDQDLDTELYSVSSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATUTTLSQDPETELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYMLENGEELNATNTTLSQDPETKLYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE KRIRCSASGDFPEPRLAMMEDGEELNAVNTTVDQDLDTELYSVSSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELMATUTTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE KRIRCSASGGFPEPRLAWMEDGEELNAVNTTVDQDLDTELYSVSSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE KRIRCSASGGFPEPRLAWMEDGEELNAVNTTVDQDLDTELYSVSSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATVTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATNTTLSQDPETKLYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATNTTLSQDPETKLYMISSELDFNWTSNHSFLCLVKYGDLTVSQSFYWQESKPT RRLICSTSGGFPRPHLYWLENGEELNATNTTLSQDPETKLYMISSELDFNVTNNRSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATUTVSQDPFTXLYMISSELDFNTTSNHSFLCLVKYGDLTVSQTFYWQESKPT RRLICSTSGGFPRPHLWLENGKELNATUTTLSQDPETKLYMISSELDFNMTSNHSFLCLVKYGDLTVSQTFYWQESKPT rrlicstsggfprphlywlengeelnatnttlsqdpetklymisseldfnvynnysivclikygelsvsqifpwskpkqe rrlicstsggfprphlywlengeelnatnttsqdpetklymisseldfnmtsn---lclvkygdlyvsqtfywqeskpt KRIRCSASGDFPEPRLAWMEDGEELNAVNTTVDQDLDTELYSVSSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATUTTVSQDPGTELYMISSELGFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE rrlicstsggfprphlywlengeelnatnttsqdpetklymisseldfnmtsnhsflclvkygdltvsqtfywqeskpt RRLICSTSGGFPRPHLYWLENGEELNATNTTVSQDPGTELYMISSELDFNVTNNHSIVCLIKYGELSVSQIFPWSKPKQE RRLICSTSGGFPRPHLYWLENGEELNATUTLSQDPETKLYMISSELDFNWTSNHSFLCLVKYGDLTVSQTFYWQESKPT RRL ICSTSGGF PR PHL WLENGEELNATNTTVSQD PGTEL YM ISSELDFNVTNNHS IVCL IKYGELSVSQIF PWSK PKQE (160)(161) (161) (161)161) (161)160) 161) (161) 161) (161) 161) 161) (161)(161)160) (161)(161)(161)(161) (159)161) (161) 161) 160) 161) 161) 161) 161) 161) 161) (161)(161) (161)158) (161)EQ:283_CD28BP_Con SEQ:196_cd28D12-5 SEQ:198_cd28F7-2 SEQ:199_cd28F8-4 SEQ:200_cd28F10-2 SEQ:204_cd28G1-9 SEQ:207_cd28H6-6 SEQ:209_cd28B4-5a SEQ:212_cd28D5-6 SEQ:213_cd28D10-4 SEQ:214_cd28E2-5star SEQ:216_cd28E8-6 SEQ:217_cd28E9-6 SEQ:219_cd28F3-5 SEQ:220_cd28F3-6 SEQ:221_cd28F11-8 SEQ:187_cd28C6-1 SEQ:188_cd28C7-3 SEQ:189_cd28C8-6 SEQ:190_cd28c9-5star SEQ:191_cd28C2-4 SEQ:192_cd28D2-3 SEQ:193_cd28D2-9 SEQ:194_cd28D8-9 SEQ:195_cd28D11-1 SEQ:197_cd28E10-6 SEQ: 201_cd28F12-5star SEQ:202_cd28G2-8 SEQ:203_cd28G1-5 SEQ:205_cd28H4-3 SEQ:206_cd28H11-3 SEQ:208_cd28E2-4 SEQ:210_cd28A2-5a SEQ:211_cd28B4-5star SEQ:215_cd28E5-2 SEQ:218_cd28F3-1

Extracellular domain (ECD)



P-PIDQLPFWVIIP---VSGALVLTAVVLYCPACRHVARWKRTRRNEETVGTERLSPIYLGSAQSRAEVPSLSX P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAOSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYRPACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRIRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAOSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP----VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP---VSGALVLAAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP---VSGALVLTVVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFLVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFRVIIP---VSGALVLTAIVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAOSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-PIDOLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAOSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAOSSG P-PIDQLPFWVIIP---VSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG P-SANQHLTWTIIIPVSAFGISVIIAVILTCLTCRNAAIRRQRRENEVEMOSCSOSP------P-SANQHLTWTIIIPVSAFGISVIIAVILTCLTCRNAAIRRQRRENEVKNQSCSQSP------P-SANQHLTWTIIIPVSAFGISVIIAVILTCLTCRNAAIRRQRRENEVEMQSCSQSP----(238) HFPDNLLPSWAITL----ISVNGIFVICCLTYCFAPRCRERRNE-RLRRESVRPV-P-PIDQLPFWVIVP---VSGALVLTAVVLYCLACRHVAR------(241) (237) (241)(241)(241) (241)(240) (241)(241)240) (241) (237)(241) (241)(241) (241)(241)(241)(241)(241)(241)(238)(241)(241)241) (241) (241) 241) (241)(241)(240) 241) (241)SEQ:050_R1_Clone_118 SEQ:067_R2_CD28BP-16 SEQ:176_cd28A4-9 SEQ:177_cd28A6-9 SEQ:181_cd28B2-8 SEQ:278_Human_B7-1 SEQ:049_R1_Clone_84 SEQ:056_R2_CD28BP-5 SEQ:057_R2_CD28BP-6 SEQ:059_R2_CD28BP-8 SEQ:060_R2_CD28BP-9 SEQ:061_R2_CD28BP-10 SEQ:064_R2_CD28BP-13 SEQ:066_R2_CD28BP-15 SEQ:068_R2_CD28BP-17 SEQ:174_cd28A12-5 SEQ:175_cd28a4-5star SEQ:179_cd28A8-4 SEQ:180_cd28A8-6 SEQ:182_cd28B4-3 SEQ:183_cd28B6-3 SEQ:184_cd28b6-6 SEQ:186_cd28c11-5 SEQ:051_R1_Clone_126 SEQ:053_R2_CD28BP-2 SEQ:055_R2_CD28BP-4 SEQ:058_R2_CD28BP-7 SEQ:062_R2_CD28BP-11 SEQ:063_R2_CD28BP-12 SEQ:065_R2_CD28BP-14 SEQ:185_cd28b8-5star SEQ:048_R1_Clone_71 SEQ:054_R2_CD28BP-3 SEQ:178_cd28A6-1 SEQ:052_R2_CD28BP-1

Fig. 2G



	241
SEQ:187_cd28C6-1	. (241) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHGARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:188_cd28C7-3	(241) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:189_cd28C8-6	(241) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:190_cd28c9-5star	 (241) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARXKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:191_cd28C2-4	. (238) P-SANQHLTWTIIIPVSAFGISVIIAVILTCLTCRNAAIRRQRRENEVEMQSCSQSP
SEQ:192_cd28D2-3	(239) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:193_cd28D2-9	(241) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:194_cd28D8-9	(241) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:195_cd28D11-1	. (240) P-PIDQLPFWVIILVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:196_cd28D12-5	. (238) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQPSG
SEQ:197_cd28E10-6	. (241) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:198_cd28F7-2	(241) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:199_cd28F8-4	. (241) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:200_cd28F10-2	(241) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:201_cd28F12-5star	. (241) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:202_cd28G2-8	: (241) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:203_cd28G1-5	. (240) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:204_cd28G1-9	(241) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:205_cd28H4-3	(241) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:206_cd28H11-3	_
SEQ:207_cd28H6-6	
SEQ:208_cd28E2-4	. (241) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:209_cd28B4-5a	_
SEQ:210_cd28A2-5a	. (241) P-SANQHLTWTIIIPVSAFGISVIIAVILTCLTCRNAAIRRQRRENEGKCKVLSVSIGTKLKFNR
SEQ:211_cd28B4-5star	. (241) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:212_cd28D5-6	. (241) P-SANQHLTWTIIIPVSAFGISVIIAVILTCLTCRNAAIRRQRRENEVEMQSCSQSP
SEQ:213_cd28D10-4	. (240) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:214_cd28E2-5star	. (241) P-SANQHLTWTIIIPVSAFGISVIIAVILTCLTCRNAAIRRQRRENEVEMQSCSQSP
SEQ:215_cd28E5-2	(241)
SEQ:216_cd28E8-6	(241) P-SANQHLTWTIIIPVSAFGISVIIAVILTCLTCRNAAIRRQRRENEVEMQSCSQSP
SEQ:217_cd28E9-6	(241) P-PIDQLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSPIYLGSAQSSG
SEQ:218_cd28F3-1	(241)
SEQ:219_cd28F3-5	_
SEQ:220_cd28F3-6	(241)
SEQ:221_cd28F11-8	(238) P-
SEQ:283_CD28BP_Con	. (241) P PIDOLPFWVIIPVSGALVLTAVVLYCLACRHVARWKRTRRNEETVGTERLSFIYLGSAQSSG

Fig. 2H

Extracellular domain (ECD) Signal sequence

(1) MGHTRRQGTSPSKCPYLNFFQLLVLAGLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMSGD

SEQ:278_Human_B7-1

SEQ:069_R1_CTLA4BP-5 SEQ:070_R1_CTLA4BP-7 SEQ:071_R1_CTLA4BP-11 SEQ:072_R1_CTLA4BP-13 SEQ:073_R1_CTLA4BP-27

(1)

(1,0,0,0)

(1)

MSHTRRQGTSPSKCPYLKFFQLLVLASLSHFCSGVIHMTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMSGD MGHTRRQGISPSKCPYLKFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMSGD MSHTRRQGISPSKCPYLNFFQLLVLASLSHFCSGVIHVTKEVKEVATLSCGLNVSVEELAQTRIYWQKEKKMVLTMMSGD

MGHTRRQGISPSKCPYLKFFQLLVLAGLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MGYTRRQGTSPSKCPYLKFFQLLVLAGLSHLCSGVIHVTNEVKEVATLSCGHNVSGEELAQTRIYWQKEKKMVLTMMYGD MSHTRRQGISPSKCPYLKFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD

MGHTRRQGISPPKCPYLNFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTPMSGD

MGHTRRQGTSPSKCPYLKFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MGYTRRQGTSPSKCPYLKFFQLLVLACLSHFCSGVIHVTREVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD

MSHTRRQGISPSKCPYLKFFQLLVLASLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKWVLTMMSGD

SEQ: 074_R2_CTLA4BP-5x2-10c SEQ: 075_R2_CTLA4BP-5x2-11d SEQ: 076_R2_CTLA4BP-5x2-12F SEQ: 077_R2_CTLA4BP-5x2-22 SEQ: 078_R2_CTLA4BP-5x2-3c SEQ: 079_R2_CTLA4BP-5x2-4c SEQ: 080_R2_CTLA4BP-5x2-4c SEQ: 081_R2_CTLA4BP-5x2-7b SEQ: 081_R2_CTLA4BP-5x2-7b

(1)

SEQ: 081_K2_CTLA4EF-5x2_8C SEQ: 082_R2_CTLA4EF-5x3-10e SEQ: 084_R2_CTLA4EF-5x3-11b SEQ: 085_R2_CTLA4EF-5x3-6f SEQ: 086_R2_CTLA4EF-5x4-11d SEQ: 086_R2_CTLA4EF-5x4-11c SEQ: 088_R2_CTLA4EF-5x4-1f SEQ: 088_R2_CTLA4EF-5x4-1f SEQ: 088_R2_CTLA4EF-5x5-2e

SEQ:090_R2_CTLA4BP-5x6-9d SEQ:091_R2_CTLA4BP-5x8-1f SEQ:092_R2_CTLA4BP-5x9-12c SEQ:222_ctla5x9d10

2_R2_CTLA4BP-5x9-12c SEQ:222_ctla5x9d10 SEQ:223_ctla5x6f6 SEQ:224_ctla5x5h12 SEQ:225_ctla5x5c10

10/39

MSHTRRQGISPSKCPYLNFFRLLVLASLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MSHTRRQGTSPSKCPYLKFFQLLVLASLSHFCSGVIHMTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMSGD MGYTRRQGISPSKCPYLKFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMSGD

(1)

(1)

MSHTRROGTSPSKCPYLKFFQLLVLASLSHFCSGVIHVTKEVKEVATLSCGLNVSVEELAQTRIYWQKEKKMVLTMMSGD

MSHTRRQGISPSKCPYLKFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMSGD MGHTRRQGISPSKCPYLNFPQLLVLAGLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIYWQKGKKMVLTMMSGD MGHTRRQGTSPSKCPYLKFFQLLVMACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MGHTRRQGISPSKCPYLKFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGLNVSVEELAQTRIHWQKEKKMVLTMMSGD MGYTRRQGISPSKCPYLKFFQLLVLASLSHFCSGVIHVTKKVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MGHTRRQGISPSKCPYLKFFQLLVLACLSHLCSGVIHVTKEVKEVATLSCGLNVSVEELAQTRIHWQKEKKMVLTMMSGD MGHTRROGISPSKCPYLKFFOLLVLAGLPHLCSGVIHVTKEVKEVATLSCGHNVSVEELAOTRIHWOKEKKMVLTMMSGD MSHTRRQGTSPSKCPYLKFFQLLVLAGLSHLCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MGHTRRQGISPSKCPYLNFPQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MGHTRRQGTSPSKCPYLNFPQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MSHTRRQGTSPSKCPYLKFPQFLVLASLSHFCSGVIHVTKEVKEVATLSCGLNVSVEELAQTR1YWQKGKKMVLTMMSGD MGYTRRQGTSPSKCPYLKFFQLLVLASLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTPIYWQKEKKMVLTMMSGD MGHTRROGTSPSKCPYLKFFQLLVLAGLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MSHTQRQGISPSKCPYLNFFQLLVLASLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMSGD

Fig. 3A

Extracellular domain (ECD) Signal sequence

1/39 MGHTRRQGTSPSKCPYLKFFQLLVLAGLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTWMSGD MGYTRRQGISPSKCPYLKFFQLLVLAGLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MSHTRRQGISPSKCPYLKFFQLLVLAGLSHFCSGVIHVTKEVKEVATLSCGHNVSAEELAQTRIYWQKEKKMVLTMMSGD MGHTRRQGISPSKCPYLKFFQLLVLAGLSHFCSGVIHVTKEVKEVATLSCGLNVSVEELAQTRIHWQKEKKMVLTMMSGD MGHTRRQGTSPSKCPYLKFFQLLVLAGLSHLCSGVIHVTKEVKEVATLSCGHNVSVEELAQTR1YWQKEKKMVLTMMSGD MSHIRRQGISPSKCPYLNFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMSGD MSHTRRQGTSPSKCPYLKFFQLLVLASLSHFCSGVIHMTKEVKEVATLSCGPNVSVEELAQTRIYWQKEKKMVLTMMSGD MSHTRRQGISSSKCPYLKFFQLLVLACLSHFCSGVIHVTKKVKEVATLSCGHNVSVEELAQTRIYWQKGKKMVLTMMSGD MGYTRRQGTSPSECPYLKFFQLLVLAGLSHFCSGVIHMTKEVKEVATLSCGLNVSVEELAQTR1HWQKEKKMVLTWMSGD MSHTRRQGISPSKCPYLNFFRLLVLASLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MSHTRRQGI SPSKCPYLKFFQLLVLASLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRI YWQKEKKMVLTMMSGD MGHTRRQGISPSKCPYLKFPQLLVLASLSHFCSGVIHVTKEVKEVATLSCGLNVSVEELAQTRIYWQKEKKMVLTMMSGD MGYTRRQGTSPSKCPYLNFFQLLVLASLSHFCSGVIHVTKEVKEVATLSCGHNVPVEELAQTRIYWOKEKKMVLTMMSGD MGHTRRQGISPSKCPYLKFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIYWQKDKKMVLTMMSGD MGHTRRQGISPSKCPYLKFFQLLVLACLSHFCSGVIYVTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLIMMSGD MGHTRRQGTSPSKCPYLKFFQLLVLAGLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MGYTRRQGTSPSKCPYLNFFQLLVLASLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MGYTRRQGISPSKCPYLKFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSDEELAQTRIHWQKEKKMVLTMMSGD MGYTRRQGISPSKCPYLKFFQLLVLAGLSHLCSGVIHVTKEVKEVATLPCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MGHTRRQGISPSKCPYLKFFQLLVLAGLSHLCSGVIHMTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMSGD MGHTRRQGISPSKCPYLKFFQLLGLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MSHTRRQGISPSKCPYLKFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD MGHTRRQGTSPSKCPYLKFFQLLVLAGLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMSGD MSHTRRQGISPSKCPYLKFFQLLVLASLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMPGD MSHTRRQGISPSKCPYLKFFQLLVLAGLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGG MGYTRRQGTSPSKCPYLKFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTR1YWQKEKKMVLTMMSGD MGHTRRQGTSPSKCPYLNFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIYWQKEKKMVLTMMSGD MGHTRRQGISPSKCPYLKFFQLLVLACLSHFCSGVIHVTKEVKEVATLSCGHNVSVEELAQTRIHWQKEKKMVLTMMSGD (1) (1) Ξ 1 $\overline{\mathbf{1}}$ 9.0 (I) 9.5 1 $\widehat{\Xi}$ (1) $\widehat{\mathbf{G}}$ (1) (1) (1) (1) (1) (1) (1) (1) £ £ SEQ:228_ctla5x3c3 SEQ:232_ctla5x2b1 ns SEQ:235_ctla2x4g9 SEQ:236_ctla2x4a6 SEQ:239_ctla2x1g8 SEQ:240_ctla2x1f10 SEQ:241_ctla2x1c9 SEQ:242_ctla2x1h12 SEQ:246_ctla2x2f1 SEQ:250_ctla5x2e12 SEQ:226_ctla5x3e8 SEQ:230_ctla5x2d7 SEQ:231_ctla5x2b7 SEQ:234_ctla5x1d7 SEQ:237_ctla2x2f3 SEQ:243_ctla2x1e2 SEQ:244_ctla2x1c4 SEQ:245_ctla2x1b12 SEQ:247_ctla5x4h1 SEQ:248_ctla5x4a1 SEQ:251_ctla2x4h11 SEQ:252_ctla2x3h2 SEQ:227_ctla5x3c4 SEQ:229_ctla5x2h11 SEQ:233_ctla5x1f1 SEQ:238_ctla2x2f12 SEQ:249_ctla5x2f3 SEQ:286_CTLA4BP_Con

MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLEYEKDAFKREHLAEVMLSVKADFPTPSISDFEIPPSNIRRI MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVMLSVKADFPTPSITDFEIPPSNIRRI MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVMLSVKADFPTPSISDFEIPPSNIRRI MNIWPEYKNRTIFDITUNLSIVILALRPSDEGTYECVVLEYEKDAFKREHLAEVTLSVKADFPTPSITDFEIPPSNIRRI

(81)

SEQ:225_ctla5x5c10

(81)

SEQ:222_ctla5x9dl0 SEQ:223_ctla5x6f6 SEQ:224_ctla5x5hl2 49/6.

12/39

(81) MNIWPEYKNRTIFDITMNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVTLSVKADFPTPSISDFEIPTSNIRRI $\texttt{MNIWPEYKNRTIFDITMNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVTLSVKADFPTPSISDFEIPTSNIRRI$ MNIWPEYKNRTIFDI TNNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVMLSVKADFPTPSISDFEIPPSNIRRI MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVMLSVKADFPTPSITDFEIPPSNIRRI MNIWPEYKNRTIFDITMNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVTLSVKADFPTPSITDFEIPPSNIKRI MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVMLSVKADFPTPSITDFEIPTSNIRRI MNIWPEYKNRTIFDITINNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVMLSVKADFPTPSISDFEIPTSNIRRI $\mathtt{MNIWPEYKNRTIFDITMNLSIVILALRPSDEGTYECVVLKYEKDAFKQEHLAEVMLSVKADFFTPSITDFEIPPSNIRRI$ MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLEYEKDAFKREHLAEVTLSVKADFPFFSISDFEIPPSNIRRI MNIWPEYKNRTIFDITNNLSIVILALRSSDEGTYECVVLKYEKDAFKREHLAEVTLSVKADFPTPSITDFEIPPSNIRRI MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLEYEKDAFKREHLAEVMLSVKADFFTPSISDFEIPTSNIRRI MNIWPEYKNRTIFDITMNLSIVILALRPSDEGTYECVVLKYDKDAFKREHLAEVTLSVKADFPTPSISDFEIPPSNIRRI MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVMLSVKADFPTPSISDFEIFTSNIRRI MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVMLSVKADFFTPSISDFEIPPSNIRRI MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVMLSVKADFPTPSITDFEIPPSNIRRI MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVILKYEKDAFKREHLAEVMLSVKADFPTPSISDFEIPTSNIRRI MNIWPEYKNRTIFDITINNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVMLSVKADFPFPSITDFEIPPSNIRRI MNIWPEYKNRTIFDITMNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVMLSVKADFPTPSITDFEIPPSNIRRI MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVTLSVKADFPTPSISDFEIPPSNIRRI MNIWPEYKNRTIFDI TUNL SIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVMLSVKADFPTPSITDFEIPPSNIRRI MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVMLSVKADFPTPSITDFEIPPSNIRRI MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVMLSVKADFPTPSITDFEIPPSNIRRI $ext{MNIWPEYKNRTIFDITUNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVMLSVKADFPTPSISDFEIPPSNIRRI$ MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLEYEKDAFKREHLAEVMLSVKADFPTPSISDFEIPPSNIRRI MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVTLSVKAGFPTPSITDFEIPPSNIRRI (81) (81) (81)(81) (81) (81) (81) (81) (81) (81) (81) (81) (81)(81) (81) (81) (81) 81) (81) (81) (81) (81) 81) SEQ:075_R2_CTLA4BP-5x2-11d SEQ:080_R2_CTLA4BP-5x2-7b SEQ:082_R2_CTLA4BP-5x3-10e SEQ:083_R2_CTLA4BP-5x3-11b SEQ:084_R2_CTLA4BP-5x3-6f SEQ:085_R2_CTLA4BP-5x4-11d SEQ:086_R2_CTLA4BP-5x4-12c SEQ:087_R2_CTLA4BP-5x4-1f SEQ:088_R2_CTLA4BP-5x5-2e SEQ:089_R2_CTLA4BP-5x5-6e SEQ: 090_R2_CTLA4BP-5x6-9d SEQ:091_R2_CTLA4BP-5x8-1f EQ: 092_R2_CTLA4BP-5x9-12c SEQ:076_R2_CTLA4BP-5X2-12F SEQ:078_R2_CTLA4BP-5x2-3c SEQ:079_R2_CTLA4BP-5x2-4c SEQ:081_R2_CTLA4BP-5x2-8c SEQ:278_Human_B7-1 SEQ:069_R1_CTLA4BP-5 SEQ:070_R1_CTLA4BP-7 SEQ:073_R1_CTLA4BP-27 SEQ:074_R2_CTLA4BP-5x2-10c SEQ:077_R2_CTLA4BP-5x2-2g SEQ:071_R1_CTLA4BP-11 SEQ:072_R1_CTLA4BP-13

Extracellular domain (ECD)

Fig. 30

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◆	160	e8 (81) MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLEYEKDAFKREHLAEVTLSVKADFPTPSISDFEIPTSNIRRI 24 (91) MATHIDEVEKDFITENIRMI SIXII XI DESPERANDEMEN EVEKDA EVERIA E	(81)	(81)	(81)	(81)	ns (81) mniwpehknrtifditungsivilalrpsdegtyecvvlkyekdafkrehlaevm.svkadfptpsisdfeippsnirri f1 (81) mniwpehknptifditungsivilalppsproponyrikopenyaevpehlaevm svxadfensoppstrefiddenidet	(81)	39 (81) MNIWPEYKNQTIFDITMNLSIVILALRPSDEGTYECVVLKYEKDAFKQEHLAEVMLSVKADFPTPSISDFEIPPSNIRRI	a6 (81) MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVMLSVKADFPTPSISDFEIPPSNIRRI	É3 (81) MNIWPEYKNRTIFDITNNLSIVILALRPSDEGT-ECVVLKYEKDAFKREHLAEVTLSVKADFPTPSISDFEIPPSNIRRI	12 (81) MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYGCVVLEYEKDAFKREHLAEVMLSVKADFPTPSISDFEIPPSNIRRI	38 (81) MNIWPEYKNRTIFDITNNLSVVILALRPSDEGTYECVVLKYEKDAFKREHLAEVTLSVKADFPTPSITDFEIPPSNIRRI	10 (81) MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVMLSVKADFPTPSITDFEIPPSNIRRI	c9 (81) MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLEYEKDAFKREHLAEVMLSVKADFPTPSITDFEIPTSNIRRI	12 (81) MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVMLSVKADFPTPSITDFEIPPSNIRRI	elminestrippitnnesivilalippernisionesivilalippertyectvilkyekdafkrehlaevtlsvkadfptpsisdfeifpsnirri	04 (81) MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVALSVKADFPTPSITDFEIPPSNIRRI	12 (81) MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVALKYEKDAFKQEHLAEVTLSVKADFPTPSISDFEIPPSNIRRI	f1 (81) MNIWPEYKNRTIFDITNNLSIVILALRLSDEGTYECVVLKYEKDAFKREHLAEVTLSVKADFPTPSISDFEIPTSNIRRI	n1 (81) MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAFKRKHLAEVMLSVKADFPTPSISDFEIPTSNIRRI	al (81) MNIWPEHKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVTLSVKADFPTPSITDFEIPTSNIRRI	f3 (81) MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLRYEKDAFKREHLAEVTLSVKADFPTPSISDFEIPTSNIRRI	12 (81) MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVTLSVKADFPTPSISDFEIPTSNIRRI	(81)	(81)	on (81) MNIWPEYKNRTIFDITNNLSIVILALRPSDEGTYECVVLKYEKDAFKREHLAEVALSVKADFPTPSISDFEIPPSNIRRI
		SEQ:226_ctla5x3e8	SEQ:228_ctla5x3c3	SEQ:229_ctla5x2h11	SEQ:230_ctla5x2d7	SEQ:231_ctla5x2l	SEQ:232_ctla5x2b1 ns	SEQ:234_ctla5x1d7	SEQ:235_ctla2x4g	SEQ:236_ctla2x4a6	SEQ:237_ctla2x2f3	SEQ:238_ctla2x2f12	SEQ:239_ctla2x1c	SEQ:240_ctla2x1f10	SEQ:241_ctla2x1c9	SEQ:242_ctla2x1h12	SEQ:243_ctla2x1e2	SEQ:244_ctla2x1c4	SEQ:245_ctla2x1b1	SEQ:246_ctla2x2f1	SEQ:247_ctla5x4h1	SEQ:248_ctla5x4a	SEQ:249_ctla5x2f3	SEQ:250_ctla5x2e12	SEQ:251_ctla2x4h11	SEQ:252_ctla2x3h2	SEQ:286_CTLA4BP_Con

Fig. 3D

I CSTSGGFP EPHLFWLENGEELNA I STTVSQDP ETEL YAVSSKLDFNMTTNHSFMCL I KYGHLRVNQTFNWNTTKQEHFP

(161)

SEQ:225_ctla5x5c10

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(161) ICSTSGGFPEPHLSWLENGEELNAINTTVSQDFGTELYAVSSKLDFNMTTNHSFMCLIKYGHLRVNQTFNWNTTKQEHFP

Extracellular domain (ECD)

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Fig. 3E

ICSTSGGFPEPHLFWLENGEELNAINTVSQDPETELYTVSSKLDFNMTANHSFVCLIKYGHLRVNQTFNWNTPKQEHFP ICSTSGGFPEPRLAMMEDGEELNAINTTASQDPETELYTVSSKLDFNMTTNRSFVCLIKYGHLRVNQTFNWNTPKQEHFP I CSTSGGFPEPHLSWLENGEELNAINTTVSQDPGTELYTVSSKLDFNMTTNHSFMCLIKYGHLRVNQTFNWNTPKQEHFP I CSTSGGFPEPHLFWLENGEELNA INTTASODPETELYAVSSKLDFNMTTNHSFMCL I KYGHLRVNOTFNWNTPKQEHFP 1CSTSGGFPEPHLSWLENGEELNAINTTVSQDPETGLYTVSSKLDFNMTTNHSFMCLIKYGHLRVNQTFNWNTPKQEHFP 1CSTSGGFPEPHLSWLENGEELNAINTTVSQDPGTELYTVSSKLDFNMTANHSFVCLIKYGHLRVNQTFNWNTPKQEHFP ICSTSGGFPEPHLSWLENGEELNAINTTVSQDPETELYTGSSKLDFNMTTNHSFWCLIKYGHLRVNQTFSWNTPRQEHFP ICSTSGGFPEPHLFWLENGEELMAINTTASQDPETELYTVSSKLDFNMTANHSFVCLIKYGHLRVNQTFNWNTPKQEHFP ICSTSGGFPEPHLFWLENGEELNAINTTASQDPETELYAVSSKLDFNMTTNHSFMCLIKYGHLRVNQTFNWNTPRQEHFP ICSTSGGFPEPRLAWMEDGEELNAISTTVSQDPGTELCTVSSKLDFNMTTNHSFMCLIRYGHLRVNQTFNWNTPKQEHFP 1CSTSGGFPEPHLFWLENGEELNA1STTVSQDPETELYAXSSKLDFNMTTNHSFMCL1KYGHLRVNQTFNWNTPKQEHFP ICSTSGGFPEPHLSWLENGEELNAINTTVSQDPGTELYTVSSKLDFNMTTNHSFMCLIKYGHLRVNOTFNWNTPKQEHFP 1CSTSGGFPEPHLSWLENGEELNAINTVSQDPETELYAVSSKLDFNMTTNHSFMCLIKYGHLRANQTFNWNTPKQEHFP ICSTSGGFPEPHLSWLENGEELNAINTTVSQDPGTELYTVSSKLDFNMTTNRSFVCLIKYGHLRVNQTFNWNTPKQEHFP ICSTSGGFPEPRLAWMEDGEELNAINTTVSQDPGTELYAVSSKLDFNMTTNHSFMCLIKYGHLRVNQTFNWNTPKQEHFP 1CSTSGGFPEPRLAWMEDGEELNA1STTASQDPETELYTVSSKLDFNMTTNHSFMCL1KYGHLRVNQTFNWNTPKQEHFP ICSTSGGFPEPHLSWLENGEELNAINTTVSQDPGTELYTVSSKLDFNMTTNHSFMCLIKYGHLRVNQTFNWNTPKQEHFP ICSTSGGFPEPHL.SWLENGEELNAINTTVSQDPGTELYTVSSKLDFWMTTNHSFWCLIKYGHLRVWQTFNWNTPKQEHFP ICSTSGGFPEPHLFWLENGEELNAINTTVSQDPGTELYAVSSKLDFNMTTNHNFMCLIKYGHLRVNQTFNWNTPKQEHFP 1CSTSGGFPEPRLAWMEDGEELNAINTVSQDPETELYTVSSKLDFNMTANHSFMCL1KYGHLRVNQTFNWNTPKQEHFP ICSTSGGFPEPHLFWLENGEELNAINTTASQDPETELYTVSSKLDFNMTTNRSFVCLIKYGHLRVNQTFNWNTPKQEHFP ICSTSGGFPEPHLFWLENGEELNAINTTASQDPETELYTVSSKLDFNMTTNRSFVCLIKYGHLRVNQTFNWNTPKQEHFP I CSTSGGFPEPHLSWLENGEELMAINTTVSQDPGTELYTVSSKLDFNMTTNRSFVCLIKYGHLRVNQTFNWNTPRQEHFP ICSTSGGFPEPHLSWLENGEELNAINTTASQDPETELYTVSSKLDFNMTTNRSFVCLIKYGHLRVNQTFNWNTPKQEHFP ICSTSGGFPEPHLSWLENGEELNAI STTVSQDPGTELYAVSSKLDFNMTTNRSFVCL IKYGHLRVNQTFNMNTTKQEHFP ICSTSGGFPEPHLSWLENGEELNAINTTVSQDPGTELYAVSSKLDFNMTTNHSFMCLIKYGHLRVNQTFNMNTPKQEHFP 1CSTPGGFPEPRLAWMEDGEELMA1STTVSQDPGTELYAVSSKLDFNMTTNHSFMCL1KYGHLRVNQTFNWNTTKQEHFP ICSTSGGFPEPHLSWLENGEELNAINTVSQDPETEL?TVSSKLDFNMTTNHSFMCLIKYGHLRVNQTFNWNTPKQEHFP (161)(161)(161)(161)(161)(161)(161)161) (161)(161)(161)(161)(161) (161)(161)(161)(161)(161)(161)(161)(160)(161)(161)(161)(161)(161)SEQ:227_ctla5x3c4 SEQ:228_ctla5x3c3 SEQ:232_ctla5x2b1 ns SEQ:235_ctla2x4g9 SEQ:236_ctla2x4a6 SEQ:238_ctla2x2f12 SEQ:239_ctla2x1g8 SEQ:242_ctla2x1h12 SEQ:243_ctla2x1e2 SEQ:244_ctla2x1c4 SEQ:245_ctla2x1b12 SEQ:246_ctla2x2f1 SEQ:250_ctla5x2e12 SEQ:252_ctla2x3h2 SEQ:226_ctla5x3e8 SEQ:229_ctla5x2h11 SEQ:230_ctla5x2d7 SEQ:231_ctla5x2b7 SEQ:233_ctla5x1f1 SEQ:234_ctla5x1d7 SEQ:237_ctla2x2f3 SEQ:240_ctla2x1f10 SEQ:241_ctla2x1c9 SEQ:247_ctla5x4h1 SEQ:248_ctla5x4a1 SEQ:249_ctla5x2f3 SEQ:251_ctla2x4h11 SEQ:286_CTLA4BP_Con

Extracellular domain (ECD)

Fig. 3F

DNLL PSWAITLISVNGIFVICCLTYCFAPRCRERRRNERLRRESVHPV



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(241) DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERRRNERLRRESVRPV DNLLPSWAITLISANGIFVICCLTYCFAPRCRERRRBERLRRESVHPV DNLLPSWAITLISANGIFVICCLAYCFAPGCRERKSNERLRRESVRPV DNLL PSWAITL I SVNGI FVICCL TYCFA PRCRERRRNERLRRESVCPV DNLL PSWAITL I SVNGIFVICCL TYCFA PRCRERRRNERLRRESVRPV DNLL PSWAITLISANGIFVICCL TYCFA PRCRERKSNERLRRESVRPV DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERRNETLRRESVRPV DNLL PSWAITLISANGIFVICCL TYCFAPRCRERKSNETLRRESVRPV DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERRNERLRRESVCPV DNLL PSWAITLISVNGIFVICCLTYCFAPRCRERRRNERLRRESVHPV DNLL PSWAITLISVNGIFVICCLTYCFAPRCRERRSNERLRRESVRPV DNLL PSWAITLISVNGIFVICCL TYRFA PRCRERKSNERLRRESVRPV DNLL PSWAITLISANGIFVICCL TYRFA PRCRERKSNETLRRESVRPV DNLL PSWAITL ISVNGIFVICCL TYCFA PRCRERRRNETLRRESVRPV DNLL PSWAITL I SVNGI FVICCL TYCFA PRCRERK SNERLRRESVR PV DNLLPSWAITLISVNGIFVICCLTHCFAPRCRERRRNERLRRESARPV DNLL PSWAITL I SANGI FVICCL TYRFA PRCRERRNERLRRESVCPV DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERRNGRLRRESVRPV DNLL PSWAITLISVNGIFVICCLTYCFA PRCRERRNERLRRESVRPV DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERRRNERLRRESVRPV DNLL PSWAITLISVNGIFVICCLTHCFA PRCRERRRNERLRRESVHPV DNLL PSWAITLISVNGIFVICCL TYCFA PRCRERR-NETLRRESVRPV DNPLPSWAITLISANGIFVICCLTYCFAPRCRERRRNETLRRESVRPV DNLLPSWAITLISVNGIFVICCLTYRFAPRCRERKSNERLRRESVRPV DNLLPSWAITLISANGIFVICCLTYCFAPRCRERRNERLRRESVRPV DNLLPSWAITLISANGIFVICCLTYCFAPRCRERKSNERLRRESVHPV DNLLPSWAITLISANGIFVICCLTHCFAPRCRERKRNERLRRESVRPV DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERKSNERLRRESVRPV (241)(241)(241)(241) (241)(241)(241) (241)(241)(241)(241)(241)(241)(241)(241)(241) (241) (241)(241)(241)241) 241) 241) 241) (241) (241) (241) (241)SEQ:082_R2_CTLA4BP-5x3-10e SEQ:083_R2_CTLA4BP-5x3-11b SEQ:085_R2_CTLA4BP-5x4-11d SEQ:088_R2_CTLA4BP-5x5-2e SEQ:089_R2_CTLA4BP-5x5-6e SEQ:091_R2_CTLA4BP-5x8-1f SEQ:278_Human_B7-1 SEQ:069_R1_CTLA4BP-5 SEQ:075_R2_CTLA4BP-5x2-11d SEQ:076_R2_CTLA4BP-5X2-12F SEQ:078_R2_CTLA4BP-5x2-3c SEQ:080_R2_CTLA4BP-5x2-7b SEQ:084_R2_CTLA4BP-5x3-6f SEQ:086_R2_CTLA4BP-5x4-12c SEQ:087_R2_CTLA4BP-5x4-1f SEQ:090_R2_CTLA4BP-5x6-9d EQ:092_R2_CTLA4BP-5x9-12c SEQ:222_ctla5x9d10 SEQ:225_ctla5x5c10 SEQ:074_R2_CTLA4BP-5x2-10c SEQ:077_R2_CTLA4BP-5x2-2g SEQ:079_R2_CTLA4BP-5x2-4c SEQ:081_R2_CTLA4BP-5x2-8c SEQ:224_ctla5x5h12 SEQ:070_R1_CTLA4BP-7 SEQ:072_R1_CTLA4BP-13 SEQ:073_R1_CTLA4BP-27 SEQ:071_R1_CTLA4BP-11

DNLL PSWAITL I SANGIFVICCLTYCFA PRCRERRNETL RRESVR PVWGTKLKFK PXIS DNLLPSWAITLISVNGIFVICCLTYCFAPGCRERRNERLRRESVCPV DNLFPSWAITLISVNGIFVICCLTYCFAPRCRERRRNERLRRESVRPV DNLL PSWAITL ISVNGIFVICCLTYCFAPRCRERKSNERLRRESVRPV DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERRRNETLRRESVRPV DNLL PSWAITLISVNGIFVICCPTYCFAPRCRERRRNERLRRESVCPV DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERRNERLRRESVCPV DNLLPSWAITLISVNGIFVICCLTHCFAPRCRERRRNERLRRESVCPV DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERRRNERLRRESVCPV DNLL PSWAITLISVNGIFVICCL TYCFAPRCRERRNERLRRESVRPV DNLL PSWAITLISANGIFVICCLTHCFAPRCRERKSNERLRRESVRPV DNLL PSWAITLISVNGIFVICCLAYCFAPRCRGRRRNERLRRESVRPV DNLLPSWAITLISVKGIFVICCLTYCFAPRWRERKSNERLRRESVRPV DNLLPSWAITLISVNGIFVICCLTHCFAPRCRERRNERLRRESVCPV DNLL PSWAITLISVNGIFVICCLTYCFAPRCRERRRNERLRRESVRPV DNLLPSWAITLISANGIFVICCLTYCFAPRCRERRRNERLRRESVRPV DNLL PSWAITL I SANGIFVICCL TYCFA PRCRERRRNERL RRESVHPV DNLL PSWAITL ISVKGIFVICCLTYCFAPRGRERKSNGRLRRESVHPV DNLL PSWAITLISVNGIFVICCLTYCFAPRCRERRNERLRRESVRPV DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERKSNERLRRESVRPV DNLLPSWAITLISVNGISVICCLTYCFAPRCRERRRNERLRRESVCPV DNLL PSWAITL I SANGIFVICCL TYCFA PRCRERK SNERL RRESVCPV DNLLPS-AITLISANGIFVICCLTYCFAPRCRERRNERLRRESIHPV DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERRNERLRRESVRPV DNLLPSWAITLISVNGIFVICCLTYCFAPRCRERRNERLRRESVRPV NNLL PSWAITLISVNGI FVICCLTYCFA PRCRERRNETL RRESVHPV DNLL PSWAITLISVNGIFVICCLTYCFAPRCRERRRNERLRRESVRPV DNLL PSWAITLISVNGIFVICCLTYCFAPRCRERRRNERLRRESVRPV (241)(241)(240)(241)241) (241)ECD SEQ:226_ctla5x3e8 SEQ:227_ctla5x3c4 SEQ:228_ctla5x3c3 SEQ:232_ctla5x2b1 ns SEQ:234_ctla5x1d7 SEQ:235_ctla2x4g9 SEQ:236_ctla2x4a6 SEQ:237_ctla2x2f3 SEQ:238_ctla2x2f12 SEQ:239_ctla2x1g8 SEQ:240_ctla2x1f10 SEQ:241_ctla2x1c9 SEQ:242_ctla2x1h12 SEQ:243_ctla2x1e2 SEQ:244_ctla2x1c4 SEQ:245_ctla2x1b12 SEQ:246_ctla2x2f1 SEQ:247_ctla5x4h1 SEQ:249_ctla5x2f3 SEQ:252_ctla2x3h2 SEQ:286_CTLA4BP_Con SEQ:229_ctla5x2h11 SEQ:230_ctla5x2d7 SEQ:231_ctla5x2b7 SEQ:250_ctla5x2e12 SEQ:251_ctla2x4h11 SEQ:233_ctla5x1f1 SEQ:248_ctla5x4a1

Fig. 3H

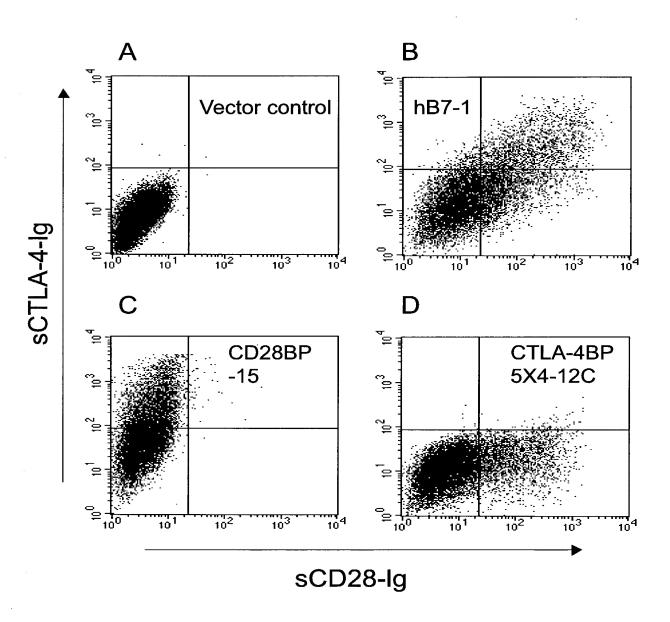
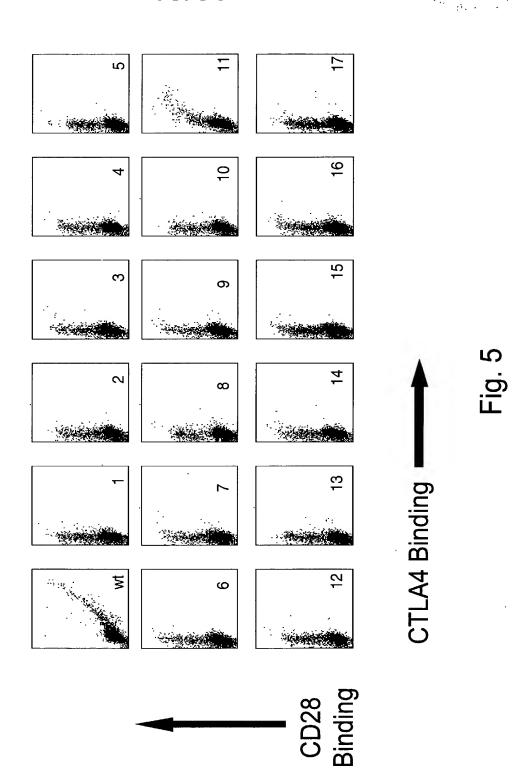
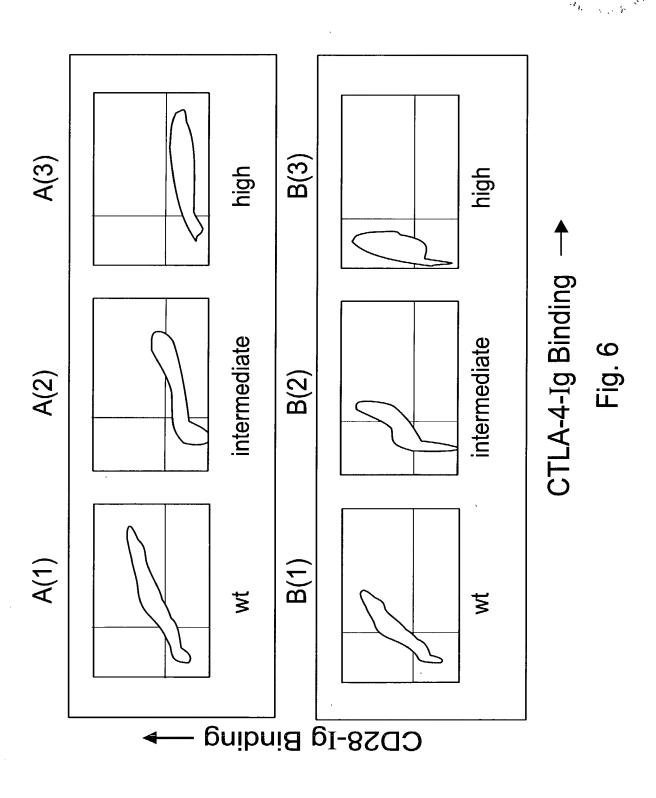


Fig. 4





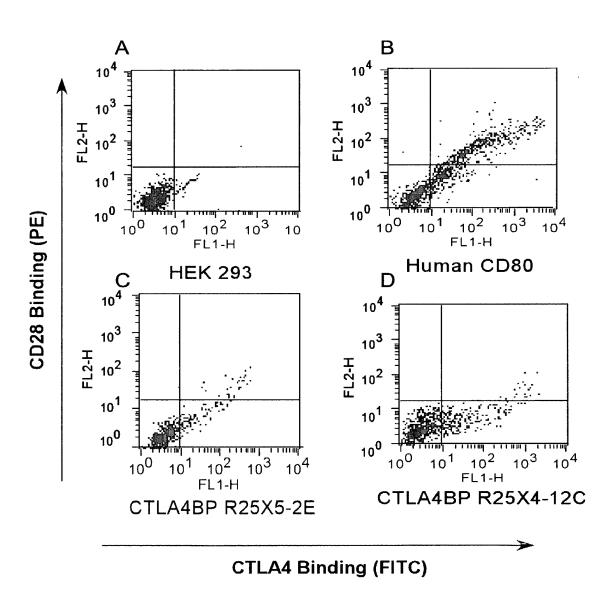


Fig. 7A-D

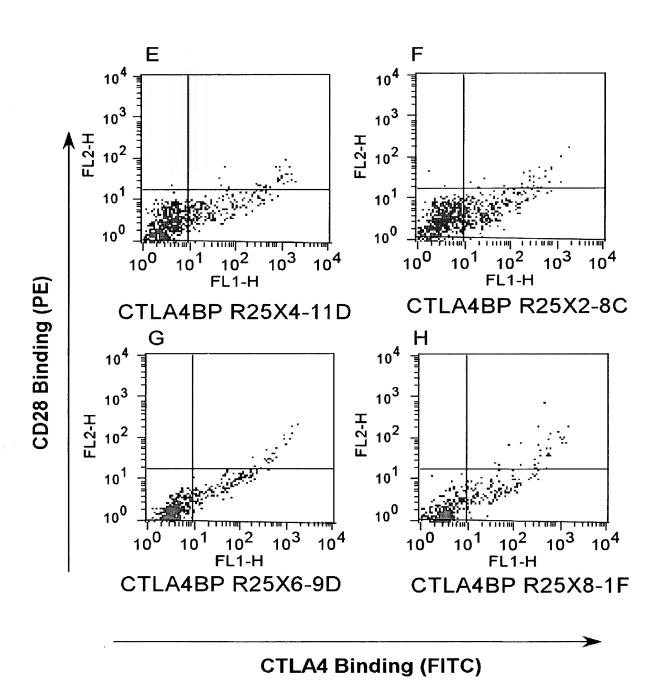
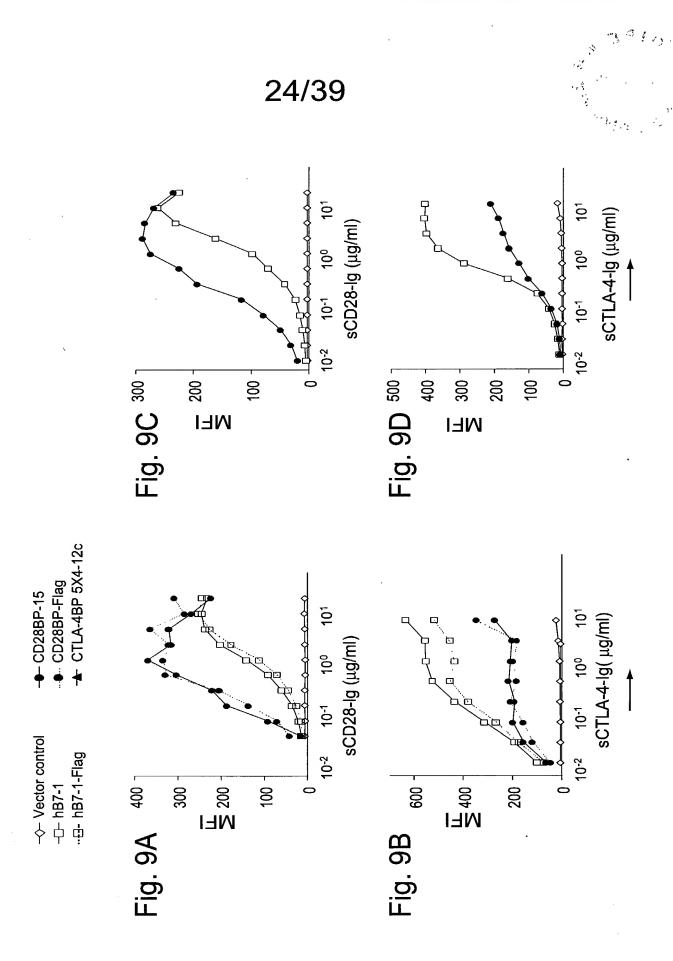
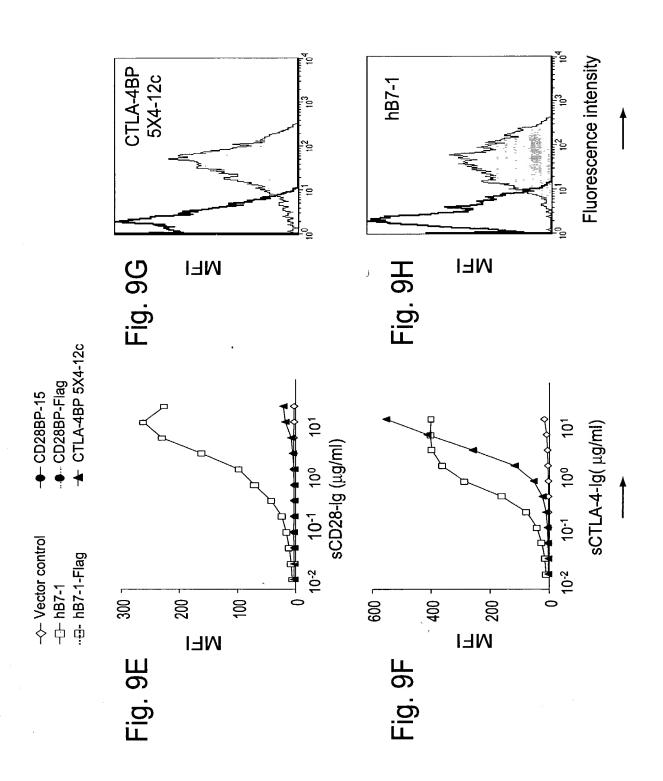


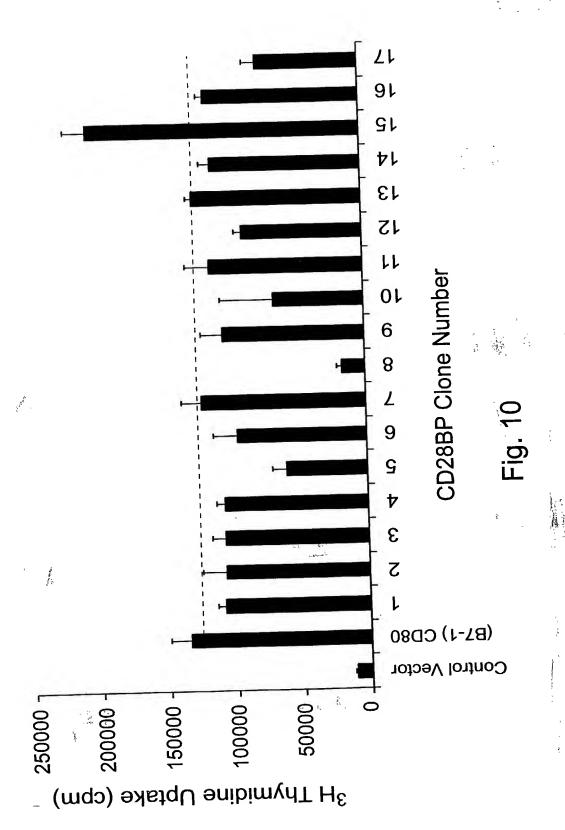
Fig. 7E-H

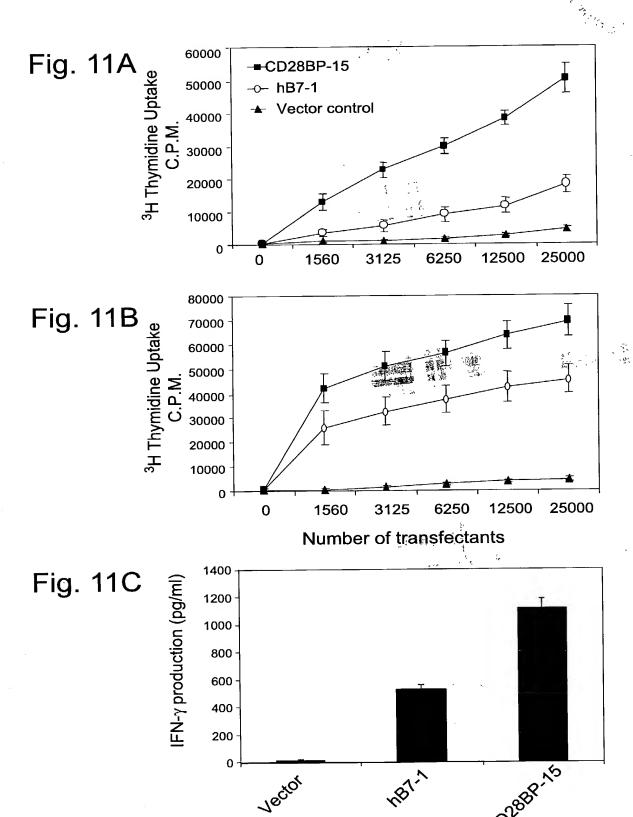
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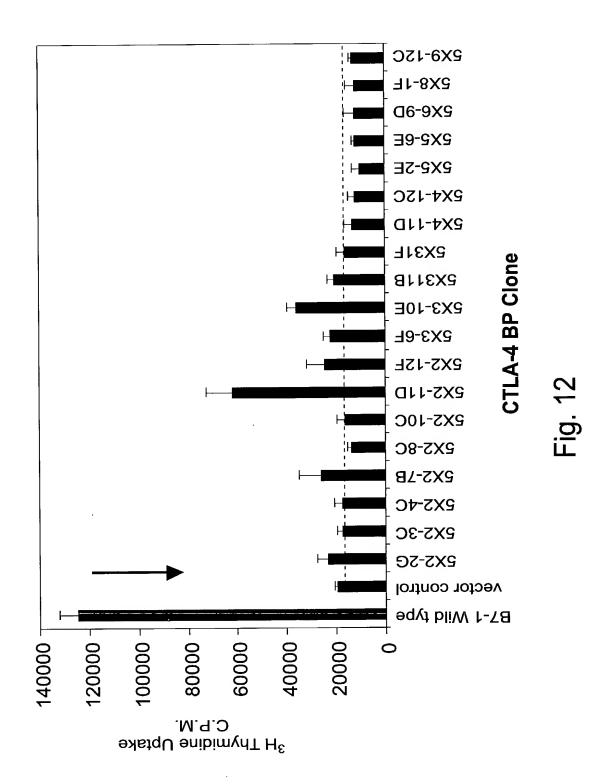














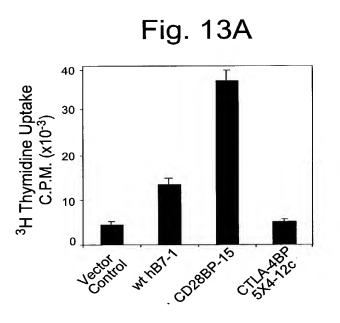
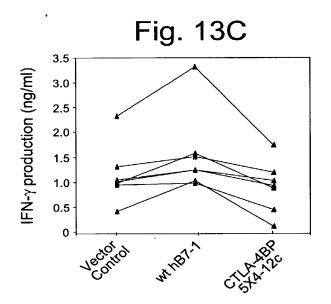
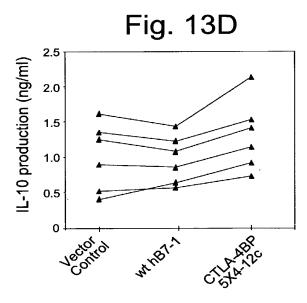


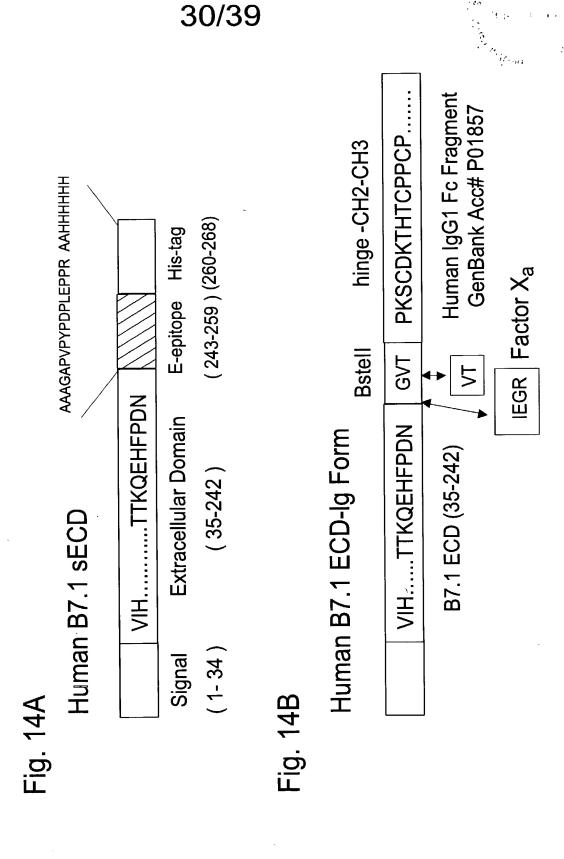
Fig. 13B

50
40
30
20
10
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Number of transfectants







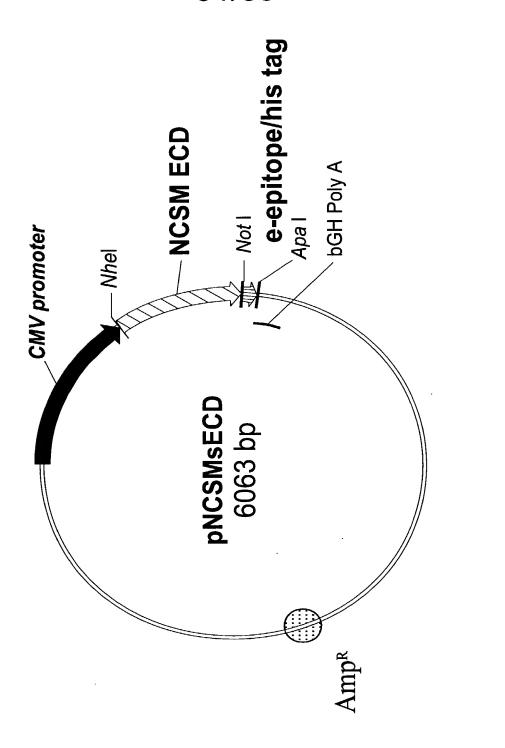


Fig. 15

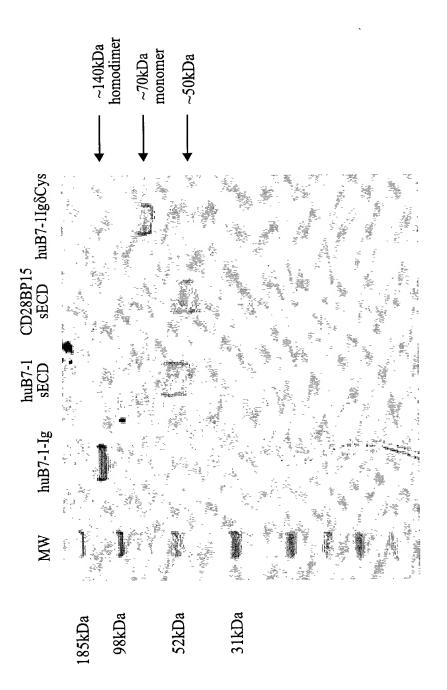


Fig. 16



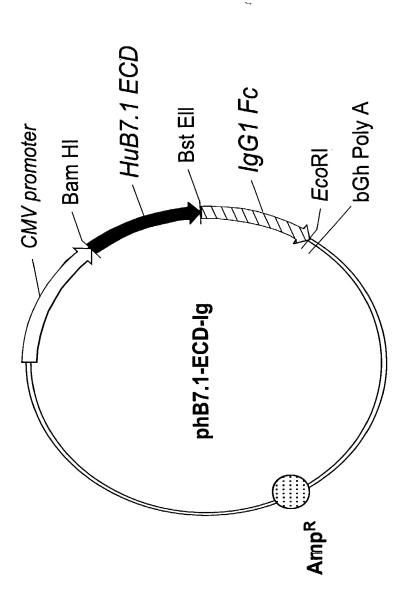


Fig. 17

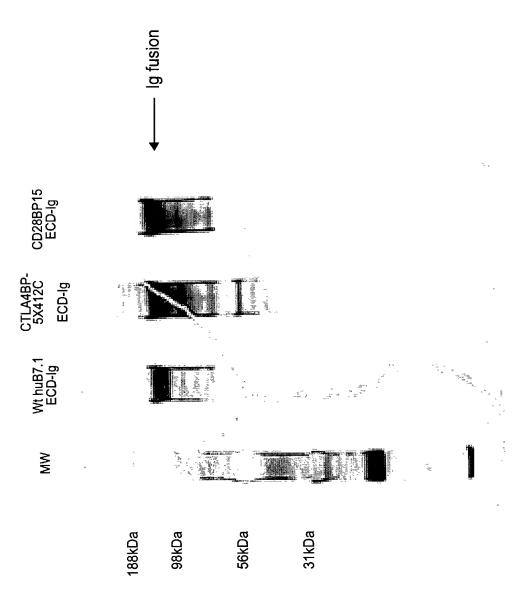


Fig. 18

Vector huB7.1-Fc WM CTLA4-5X2-8C ECD-Fc CTLA-5X4-11D ECD-Fc CTLA4-5X4-12C ECD-Fc CDS8Bb-8 ECD-Lc CD58Bb-11 ECD-Ec CD58Bb-12 ECD-Lc CTLA-45X5-2E ECD-Fc CTLA-5X6-9D ECD-Fc CTLA4-5X8-1F ECD-Fc

Expression of CTLA-4BP-Ig and CD28BP-Ig Proteins

Fig. 19

Fig. 20A

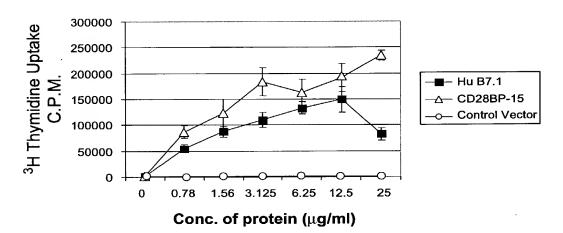
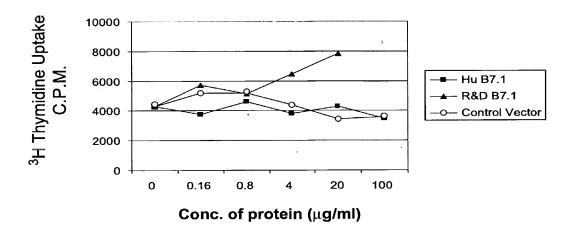
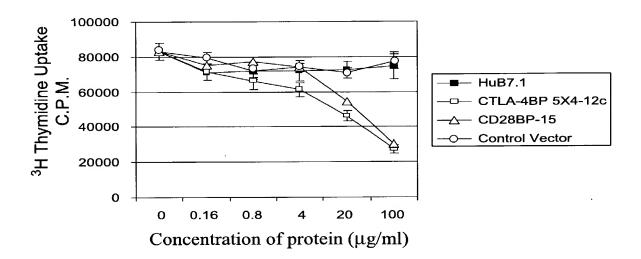


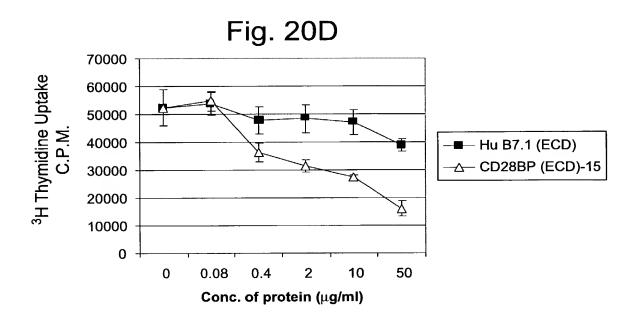
Fig. 20B



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Fig. 20C





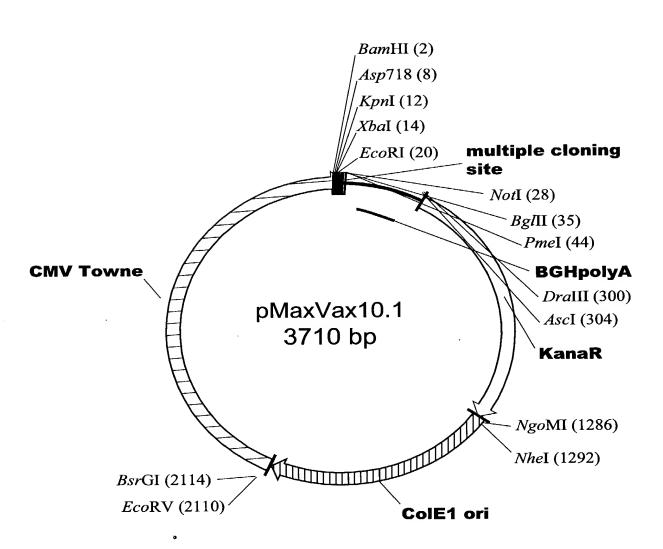


Fig. 21

Fig. 22A

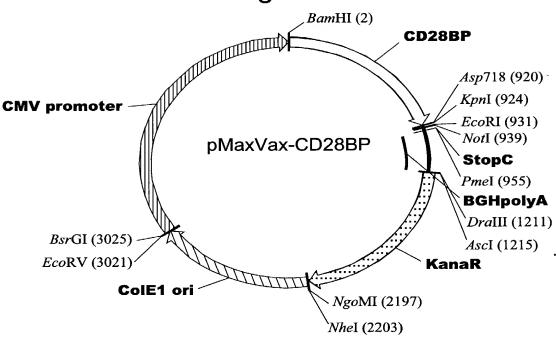


Fig. 22B

